

**Amendments to the Specification:**

Please replace the paragraph beginning at page 5, line 32, with the following rewritten paragraph:

--Figure 4 shows the ligand binding fingerprint of CCX CKR. Figure 4 Figure 4A: Definition of CCX CKR protein binding activity, as indicated by using  $^{125}\text{I}$ -ELC against a comprehensive array of viral, human, and murine chemokines in binding competition. The percent inhibition of specific binding is shown as a bar graph to emphasize that chemokines can be classed in categories as potential "high" affinity (solid bars), potential "moderate to low" affinity (hatched bars), or "no" affinity (open bars). The results are means of three determinations, the SEM in all cases is < 20%; error bars are omitted for clarity. Since intra-assay experimental error was  $\pm \sim 20\%$ , determinations within this range to the left or right of the 0% meridian are not likely to be statistically significant. Figure 4B: Rank order of high affinity CCX CKR ligand binding. Multipoint determination reflecting the competition of unlabeled chemokines against  $^{125}\text{I}$ -ELC binding to CCX CKR. Representative result of equilibrium binding using cold (unlabeled) ELC, SLC, TECK, BLC, and vMIP-II, with calculated IC50s compared in the table at bottom.--